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TECH CENTER 1600/2900

SUBSTITUTE SEQUENCE LISTING

<110> Bonnie L. Bassler
 Brendan N. Lilley

<120> LUXO-SIGMA54 INTERACTIONS AND METHODS OF
 USE

<130> PUNIV.002A

<140> 09/853,257
<141> 2001-05-10

<150> 60/202,999
<151> 2000-05-10

<160> 19

<170> FastSEQ for Windows Version 4.0

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<212> DNA
<213> Vibrio harveyi

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<213> Vibrio harveyi
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 Pro
 Ser
 Leu
 Gln
 Leu
 Lys
 Leu
 Gln
 Gln
 Leu
 Ala
 Met
 Thr
 10
 Leu
 Gln
 Leu
 Ala
 Ile
 Asp
 Leu
 Leu
 Gln
 Leu
 Ser
 Thr
 Leu
 Asp
 Asp
 Eu
 Leu
 Gln
 Leu
 Asp
 Bro
 Leu
 Asp
 Asp
 Ser
 Asp
 Asp
 Leu
 Asp
 Asp
 Ser
 Asp
 Asp
 Leu
 Asp
 Asp

100 · .105 Asn Thr Gly Ser Thr Gly Leu Ala Leu Asp Asp Asp Met Pro Val Tyr 120 Gln Gly Glu Thr Thr Glu Ser Leu His Asp Tyr Leu Met Trp Gln Leu 135 Asp Leu Thr Pro Phe Ser Glu Thr Asp Arg Thr Ile Ala Leu Ala Ile 155 150 Ile Asp Ala Val Asp Asp Tyr Gly Tyr Leu Thr Leu Ser Pro Glu Glu 165 170 Ile His Glu Ser Phe Asp Asn Glu Glu Val Glu Leu Asp Glu Val Glu 185 180 Ala Val Arg Lys Arg Ile Gln Gln Phe Asp Pro Leu Gly Val Ala Ser 200 Arg Asn Leu Gln Glu Cys Leu Leu Gln Leu Ala Thr Phe Pro Glu 215 Asp Thr Pro Trp Leu Ala Glu Ala Lys Met Val Leu Ser Asp His Ile 230 235 Asp His Leu Gly Asn Arg Asp Tyr Lys Leu Val Ile Lys Glu Ala Lys 250 Leu Lys Glu Ala Asp Leu Arg Glu Val Leu Lys Leu Ile Gln Gln Leu 260 265 Asp Pro Arg Pro Gly Ser Arg Ile Thr Pro Asp Asp Thr Glu Tyr Val 280 Ile Pro Asp Val Ser Val Phe Lys Asp His Gly Lys Trp Thr Val Ser 295 300 Ile Asn Pro Asp Ser Ile Pro Lys Leu Lys Val Asn Gln Gln Tyr Ala 310 315 Gln Leu Gly Lys Gly Asn Ser Ala Asp Ser Gln Tyr Ile Arg Ser Asn 325 330 Leu Gln Glu Ala Lys Trp Leu Ile Lys Ser Leu Glu Ser Arg Asn Glu 340 345 Thr Leu Leu Lys Val Ala Arg Cys Ile Val Glu His Gln Gln Asp Phe 360 Phe Glu Tyr Gly Glu Glu Ala Met Lys Pro Met Val Leu Asn Asp Val 375 380 Ala Leu Asp Val Asp Met His Glu Ser Thr Ile Ser Arg Val Thr Thr 390 395 Gln Lys Phe Met His Thr Pro Arg Gly Ile Phe Glu Leu Lys Tyr Phe 405 410 Phe Ser Ser His Val Ser Thr Asp Asn Gly Gly Glu Cys Ser Ser Thr 420 425 Ala Ile Arg Ala Leu Ile Lys Lys Leu Val Ala Ala Glu Asn Thr Ala 440 445 Lys Pro Leu Ser Asp Ser Lys Ile Ala Ala Leu Leu Ala Asp Gln Gly 455 460 Ile Gln Val Ala Arg Arg Thr Ile Ala Lys Tyr Arg Glu Ser Leu Gly 470 475 Ile Ala Pro Ser Ser Gln Arg Lys Arg Leu Leu 485

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<213> Vibrio harveyi

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ttggactcca accegetact ggaagttgaa gaaggecacg atgageetca ageaaatggt 180
gaagacaaat cagcgtctga atctgctgat aaaagtgcga acgaagctaa cgatgcctca 240
gaacccgacc ttccagatag ctcagacgtg attgaaaaat ctgaaatcag ctctgagcta 300
gaaattgata ccacttggga tgacgtatat agcgcaaaca cgggcagcac aggcctagcg 360
ctggatgatg acatgcccgt ctaccaaggt gagaccactg aatctttgca tgattacctt 420
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<213> Artificial Sequence
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<221> VARIANT
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<223> Xaa = Trp or Phe
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<213> Artificial Sequence
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<223> portion of consensus sequence of sigma-54 domains
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<223> upstream primer used to amplify rpoN gene
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<212> DNA
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<223> downstream primer used to amplify rpoN gene
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                                    10
Ala Ala Ser Ser Lys Ala Ser Ile Phe Ile Thr Gly Glu Ser Gly Thr
                                25
Gly Lys Glu Val Cys Ala Glu Ala Ile His Ala Ala Ser Lys Arg Gly
                            40
Asp Lys Pro Phe Ile Ala Ile Asn Cys Ala Ala Ile Pro Lys Asp Leu
                        55
                                            60
Ile Glu Ser Glu Leu Phe Gly His Val Lys Gly Ala Phe Thr Gly Ala
                    70
Ala Asn Asp Arg Gln Gly Ala Ala Glu Leu Ala Asp Gly Gly Thr Leu
                85
                                    90
Phe Leu Asp Glu Leu Cys Glu Met Asp Leu Asp Leu Gln Thr Lys Leu
            100
                                105
Leu Arg Phe Ile Gln Thr Gly Thr Phe Gln Lys Val Gly Ser Ser Lys
                            120
                                                125
Met Lys Ser Val Asp Val Arg Phe Val Cys Ala Thr Asn Arg Asp Pro
                        135
   130
                                            140
Trp Lys Glu Val Gln Glu Gly Arg Phe Arg Glu Asp Leu Tyr Tyr Arg
                   150
Leu Tyr Val Ile Pro Leu His Leu Pro Pro Leu Arg Glu Arg Gly Lys
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170
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Asp Val Ile Glu Ile Ala Tyr Ser Leu Leu Gly Tyr Met Ser His Glu
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                               185
Glu Gly Lys Ser Phe Val Arg Phe Ala Gln Asp Val Ile Glu Arg Phe
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Asn Ser Tyr Glu Trp Pro Gly Asn Val Arg Gln Leu Gln Asn
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                                25
Gly Lys Glu Leu Val Ala His Ala Leu His Arg His Ser Pro Arg Ala
                            40
Lys Ala Pro Phe Ile Ala Leu Asn Met Ala Ala Ile Pro Lys Asp Leu
                        55
Ile Glu Ser Glu Leu Phe Gly His Glu Lys Gly Ala Phe Thr Gly Ala
                                        75
Asn Thr Ile Arg Gln Gly Arg Phe Glu Gln Ala Asp Gly Gly Thr Leu
                85
                                    90
Phe Leu Asp Glu Ile Gly Asp Met Pro Leu Asp Val Gln Thr Arg Leu
           100
                               105
Leu Arg Val Leu Ala Asp Gly Gln Phe Tyr Arg Val Gly Gly Tyr Ala
                           120
Pro Val Lys Val Asp Val Arg Ile Ile Ala Ala Thr His Gln Asn Leu
                        135
Glu Arg Arg Val Gln Glu Gly Lys Phe Arg Glu Asp Leu Phe His Arg
                    150
                                        155
Leu Asn Val Ile Arg Ile His Leu Pro Pro Leu Arg Glu Arg Arg Glu
               165
                                    170
Asp Ile Pro Arg Leu Ala Arg His Phe Leu Gln Val Ala Ala Arg Glu
                               185
Leu Gly Val Glu Ala Lys Leu Leu His Pro Glu Thr Glu Thr Ala Leu
                           200
Thr Arg Leu Ala Trp Pro Gly Asn Val Arg Gln Leu Glu Asn
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Val Ser Arg Trp Asp Thr Thr Val Leu Val Arg Gly Glu Ser Gly Thr
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Gly Lys Glu Leu Ile Ala Asn Ala Ile His His Asn Ser Pro Arg Ala
                            40
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Ala Ala Phe Val Lys Phe Asn Cys Ala Ala Leu Pro Asp Asn Leu Leu Glu Ser Glu Leu Phe Gly His Glu Lys Gly Ala Phe Thr Gly Ala Val Arg Gln Arg Lys Gly Arg Phe Glu Leu Ala Asp Gly Gly Thr Leu 90 Phe Leu Asp Glu Ile Gly Glu Ser Ser Ala Ser Phe Gln Ala Lys Leu 105 Leu Arg Ile Leu Gln Glu Gly Glu Met Glu Arg Val Gly Gly Asp Glu 120 Thr Leu Arg Val Asn Val Arg Ile Ile Ala Ala Thr Asn Arg His Leu 135 Glu Glu Glu Val Arg Leu Gly His Phe Arg Glu Asp Leu Tyr Tyr Arg 150 155 Leu Asn Val Met Pro Ile Ala Leu Pro Pro Leu Arg Glu Arg Gln Glu 165 170 Asp Ile Ala Glu Leu Ala His Phe Leu Val Arg Lys Ile Ala His Ser 180 185 Gln Gly Arg Thr Leu Arg Ile Ser Asp Gly Ala Ile Arg Leu Leu Met 200 Glu Tyr Ser Trp Pro Gly Asn Val Arg Glu Leu Glu Asn 215

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                                25
Lys Glu Leu Val Ala Arg Gly Leu His Ala Ser Ser Ala Arg Ser Glu
                            40
Lys Pro Leu Val Thr Leu Asn Cys Ala Ala Leu Asn Glu Ser Leu Leu
Glu Ser Glu Leu Phe Gly His Glu Lys Gly Ala Phe Thr Gly Ala Asp
                                        75
Lys Arg Arg Glu Gly Pro Phe Val Glu Ala Asp Gly Gly Thr Cys Leu
                                    90
Asp Glu Ile Gly Asp Ile Ser Pro Met Met Gln Val Arg Leu Leu Arg
           100
                               105
Ala Ile Gln Glu Arg Glu Val Gln Arg Val Gly Ser Asn Gln Ile Ile
                            120
                                                125
Ser Val Asp Val Arg Leu Ile Ala Ala Thr His Arg Asp Leu Ala Ala
                       135
Glu Val Asn Ala Gly Arg Phe Arg Gln Asp Leu Tyr Tyr Arg Leu Asn
                   150
                                        155
Val Val Ala Ile Glu Val Pro Ser Leu Arg Gln Arg Arg Glu Asp Ile
                165
                                    170
Pro Leu Leu Ala Gly His Phe Leu Gln Arg Phe Ala Glu Arg Asn Arg
                                185
                                                    190
Arg Gly Lys Arg Phe Tyr Ala Pro Gly Leu Asp Leu Leu Ile His Tyr
                           200
Asp Trp Pro Gly Asn Ile Arg Glu Leu Glu Asn
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<213> C. crescentus
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Val Ala Pro Ser Glu Ala Ser Ile Leu Ile Thr Gly Glu Ser Gly Ser
           20
Gly Lys Glu Val Met Ala Arg Tyr Val His Gly Lys Ser Arg Arg Ala
                            40
Lys Ala Pro Phe Ile Ser Val Asn Cys Ala Ala 1le Pro Glu Asn Leu
                        55
Leu Clu Ser Glu Leu Phe Gly His Glu Lys Gly Ala Phe Thr Gly Ala
                    7.0
                                        75
Met Ala Arg Arg Ile Gly Lys Phe Glu Glu Ala Asp Gly Gly Thr Leu
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Leu Leu Asp Glu Ile Ser Glu Met Asp Val Arg Leu Gln Ala Lys Leu . 105 · 105 Leu Arg Ala Ile Gln Glu Arg Glu Ile Asp Arg Val Gly Ser Lys 120 125 Pro Val Lys Val Asn Ile Arg Ile Leu Ala Thr Ser Asn Arg Asp Leu 135 140 Ala Gln Ala Val Lys Asp Gly Thr Phe Arg Glu Asp Leu Leu Tyr Arg 150 155 Leu Asn Val Val Asn Leu Arg Leu Pro Pro Leu Arg Glu Arg Pro Ala 165 170 Asp Val Ile Ser Leu Cys Glu Phe Phe Val Lys Lys Tyr Ser Ala Ala 180 185 Asn Gly Ile Glu Glu Lys Pro Ile Ser Ala Glu Ala Lys Arg Arg Leu 200 Ile Ala His Arg Trp Pro Gly Asn Val Arg Glu Leu Glu Asn 215 <210> 16 <211> 27 <212> PRT <213> V. harveyi <400> 16 Gly Asn Ile Pro Arg Ala Ala Gly Tyr Leu Asp Val Ser Pro Ser Thr 5 Ile Tyr Arg Lys Leu Gln Ala Trp Asn Ser Lys 20 <210> 17 <211> 27 <212> PRT <213> S. typhimurium <400> 17 Gly His Lys Gln Glu Ala Ala Arg Leu Leu Gly Trp Gly Arg Asn Thr Leu Thr Arg Lys Leu Lys Glu Leu Gly Met Glu 20 <210> 18 <211> 23 <212> PRT <213> E. coli <400> 18 Gry Ash Lys Thr Glu Ala Ala Arg Gln Leu Gly Ilc Thr Arg Lys Thr ì 5 Leu Leu Ala Lys Leu Ser Arg 20 <210> 19 <211> 27 <212> PRT

<213> C. crescentus . . .

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5 10

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